

Figure 1
T. reesei egl8 nucleotide sequence (coding and non-coding)

MRATSLIAAA LAVAGDALAG KIKYLGVAIP GIDFGCDDIG SCPTDTSSVP LL₅YKGGDG₅ GQMKHFAEDD
10 20 30 40 50 60 70

GLNVFRISAT WQFVLNNNTVD GKLDLWNWGS YNKVVVNACLE TGAYCMIDMH NFARYNGGII GQGGVSDDIF
80 90 100 110 120 130 140

VDLWVQIAKY YEDNDKIIIFG LMNEPHDLDI EIWAQTCQKV VTAIRKAGAT SQMILLPGTN FASVETYVST
150 160 170 180 190 200 210

GSAEALGKIT NPDGSTDLLY FDVHKYLDIN NSGSHAECTT DNVDAFNDFA DWLRQNKRQA IISETGASME
220 230 240 250 260 270 280

PSCMTIAFCAQ NKAISENNSDV YIGFVGWGAG SFDTSYIILTL TPLGKPGNYT DNKLIMNECIL DQFTLDEKRYR
290 300 310 320 330 340 350

PTPTSISTAA EETATATATS DGDAPSTTKP IFRREETASPT PNAVTKPSPD TSDSSDDDKD SAASMSAQGL
360 370 380 390 400 410 420

TGTVLFTVAA LGYMLVAF
430 438

Predicted *T. reesei* EG VIII amino acid sequence

Figure 2